

# CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/890,779

CRF Processing Date: 4/30/02

Edited by: DC

Verified by: DC (STIC staff) # 9

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: ENTERED
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95



PCT09

## RAW SEQUENCE LISTING

DATE: 04/30/2002

PATENT APPLICATION: US/09/890,779

TIME: 15:48:33

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\04302002\I890779.raw

4 <110> APPLICANT: WOLFGANG WERR  
 6 <120> TITLE OF INVENTION: A METHOD FOR INHIBITING THE EXPRESSION  
 7 OF TARGET GENES IN PLANTS  
 10 <130> FILE REFERENCE: A34537-PCT-USA  
 12 <140> CURRENT APPLICATION NUMBER: 09/890,779  
 13 <141> CURRENT FILING DATE: 2001-08-06  
 15 <150> PRIOR APPLICATION NUMBER: PCT/EPOO/01524  
 16 <151> PRIOR FILING DATE: 2000-02-09  
 18 <150> PRIOR APPLICATION NUMBER: EP99420030.1  
 19 <151> PRIOR FILING DATE: 1999-02-09  
 21 <160> NUMBER OF SEQ ID NOS: 2  
 23 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
 25 <210> SEQ ID NO: 1  
 26 <211> LENGTH: 6  
 27 <212> TYPE: DNA  
 28 <213> ORGANISM: BACTERIA  
 30 <400> SEQUENCE: 1  
 31 gaattc 6  
 33 <210> SEQ ID NO: 2  
 34 <211> LENGTH: 8  
 35 <212> TYPE: DNA  
 36 <213> ORGANISM: BACTERIA  
 38 <400> SEQUENCE: 2  
 39 gatctcga 8

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/890,779

DATE: 04/30/2002

TIME: 15:48:34

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\04302002\I890779.raw



**Does Not Comply**  
**Corrected Diskette Needed**

PCT09

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/890,779

DATE: 04/25/2002

TIME: 12:01:12

Input Set : A:\34537 sequence listing.txt

Output Set: N:\CRF3\04252002\I890779.raw

4 <110> APPLICANT: WOLFGANG WERR  
 6 <120> TITLE OF INVENTION: A METHOD FOR INHIBITING THE EXPRESSION  
 7 OF TARGET GENES IN PLANTS  
 10 <130> FILE REFERENCE: A34537-PCT-USA  
 12 <140> CURRENT APPLICATION NUMBER: 09/890,779  
 13 <141> CURRENT FILING DATE: 2001-08-06  
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 16 <151> PRIOR FILING DATE: 2000-02-09  
 18 <150> PRIOR APPLICATION NUMBER: EP99420030.1  
 19 <151> PRIOR FILING DATE: 1999-02-09  
 21 <160> NUMBER OF SEQ ID NOS: 2  
 23 <170> SOFTWARE: FastSEQ for Windows Version 4.0

## ERRORED SEQUENCES

33 <210> SEQ ID NO: 2  
 34 <211> LENGTH: 8  
 35 <212> TYPE: DNA  
 36 <213> ORGANISM: BACTERIA  
 38 <400> SEQUENCE: 2  
 39 gatctcga  
 E--> 46 ny02:355030.1 —delete

8

VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/09/890,779

DATE: 04/25/2002

TIME: 12:01:13

Input Set : A:\34537 sequence listing.txt

Output Set: N:\CRF3\04252002\I890779.raw

Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.

Use of <220> to <223> is MANDATORY if n's or Xaa's are present.

in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:2; N Pos. 9

VERIFICATION SUMMARY

DATE: 04/25/2002

PATENT APPLICATION: US/09/890,779

TIME: 12:01:13

Input Set : A:\34537 sequence listing.txt

Output Set: N:\CRF3\04252002\I890779.raw

L:46 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:2  
L:46 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:12 SEQ:2  
L:46 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:10  
L:46 M:112 C: (48) String data converted to lower case,  
L:46 M:252 E: No. of Seq. differs, <211> LENGTH:Input:8 Found:12 SEQ:2